

003121-2266E/50

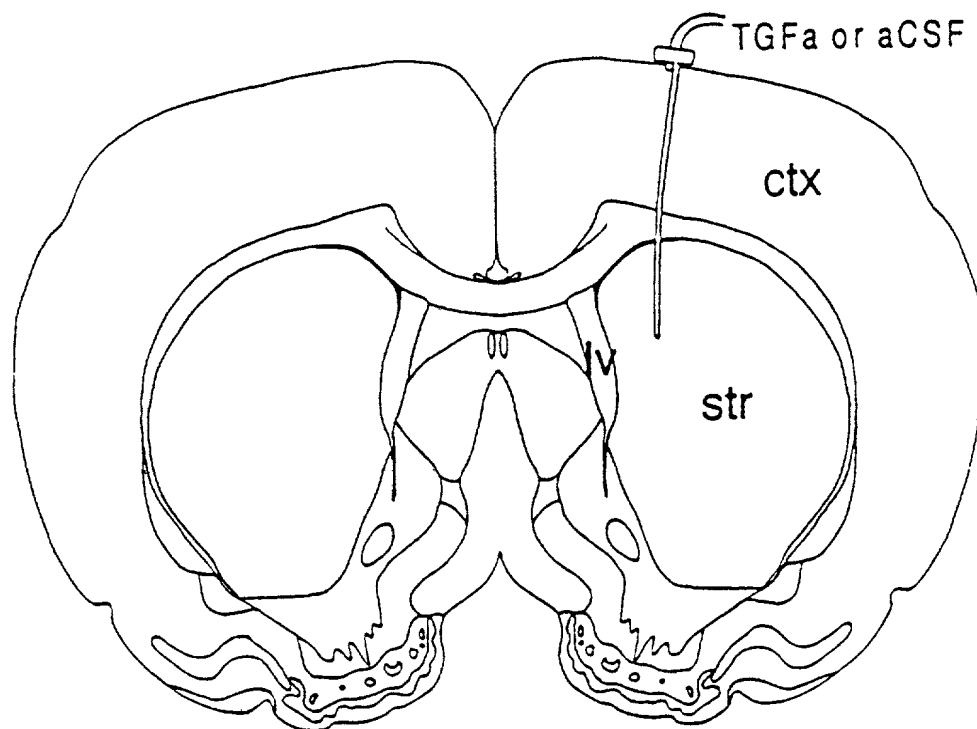
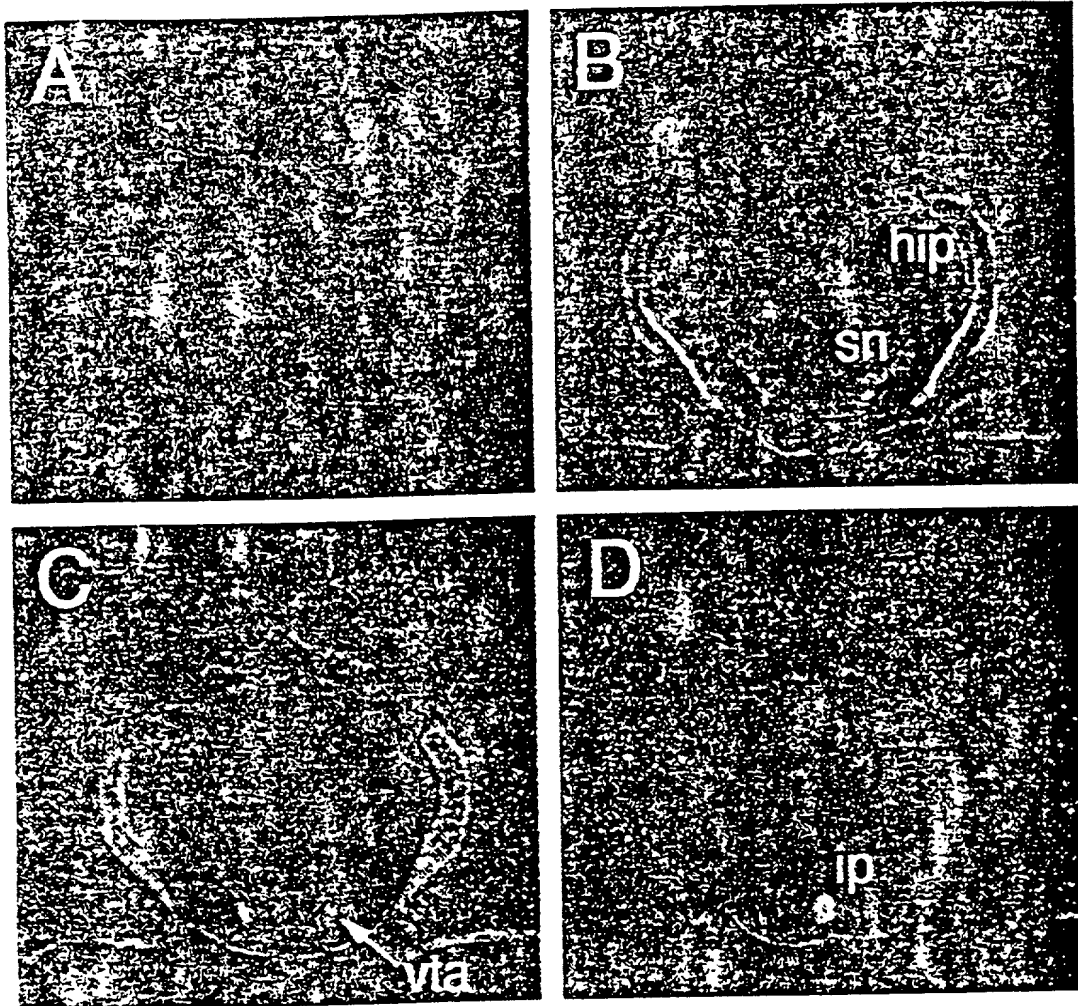


Fig. 1



Figs. 2A-2D

008T2T"EE66E/60

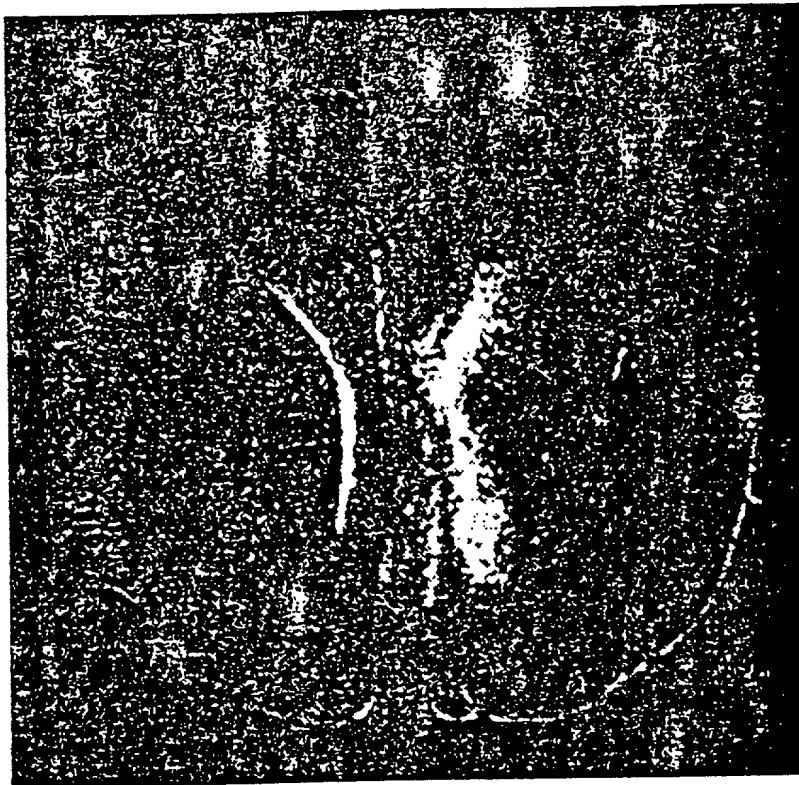


Fig. 3

008TET" E366E/60



Fig. 4

TGF α mRNA Hybridization Densities in the Striatum

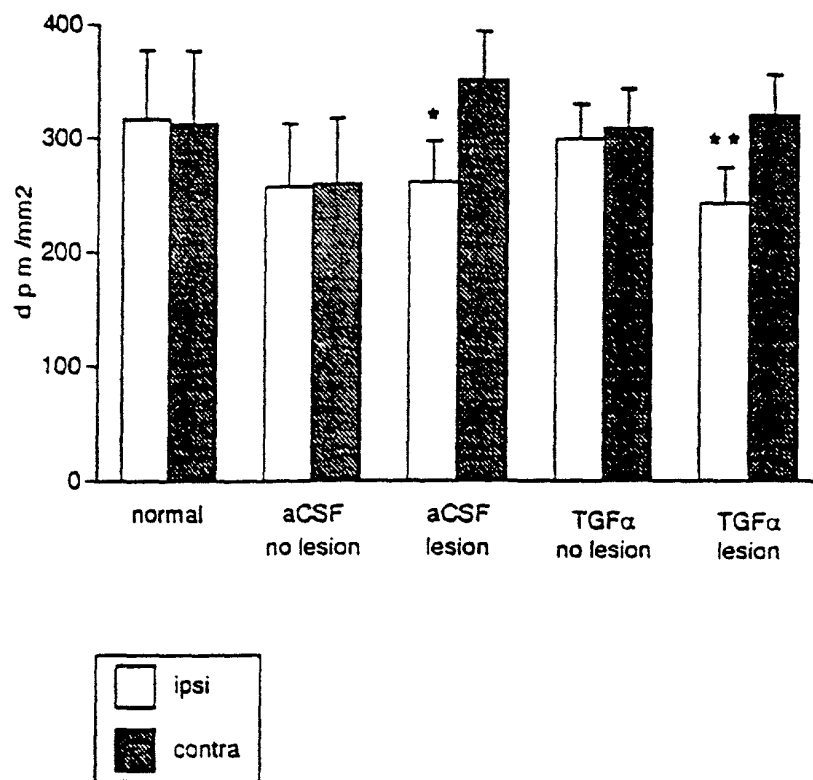


Fig. 5

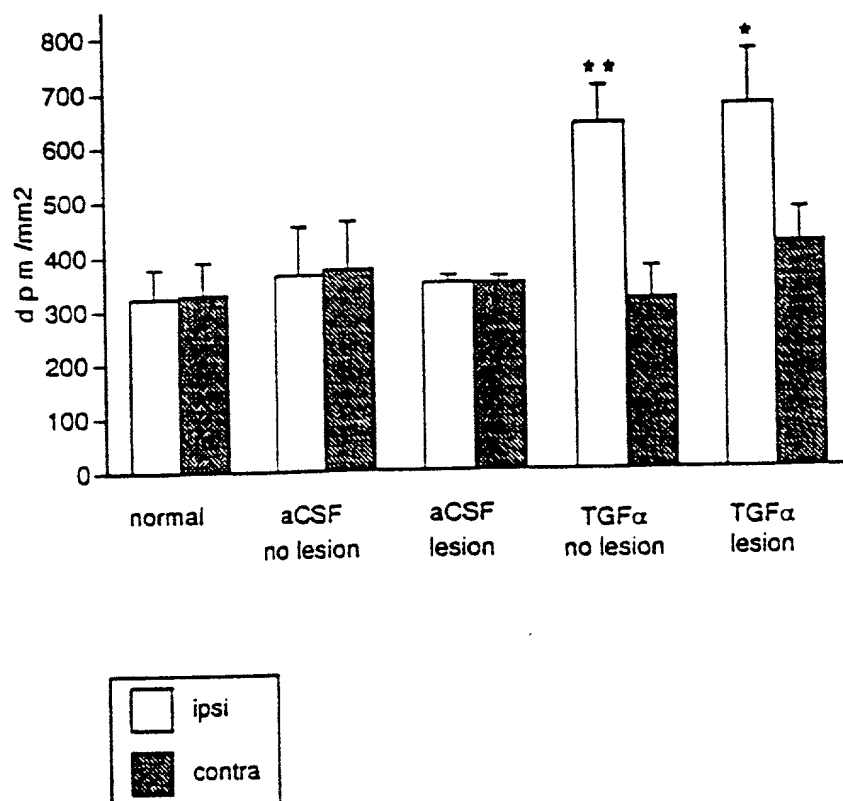
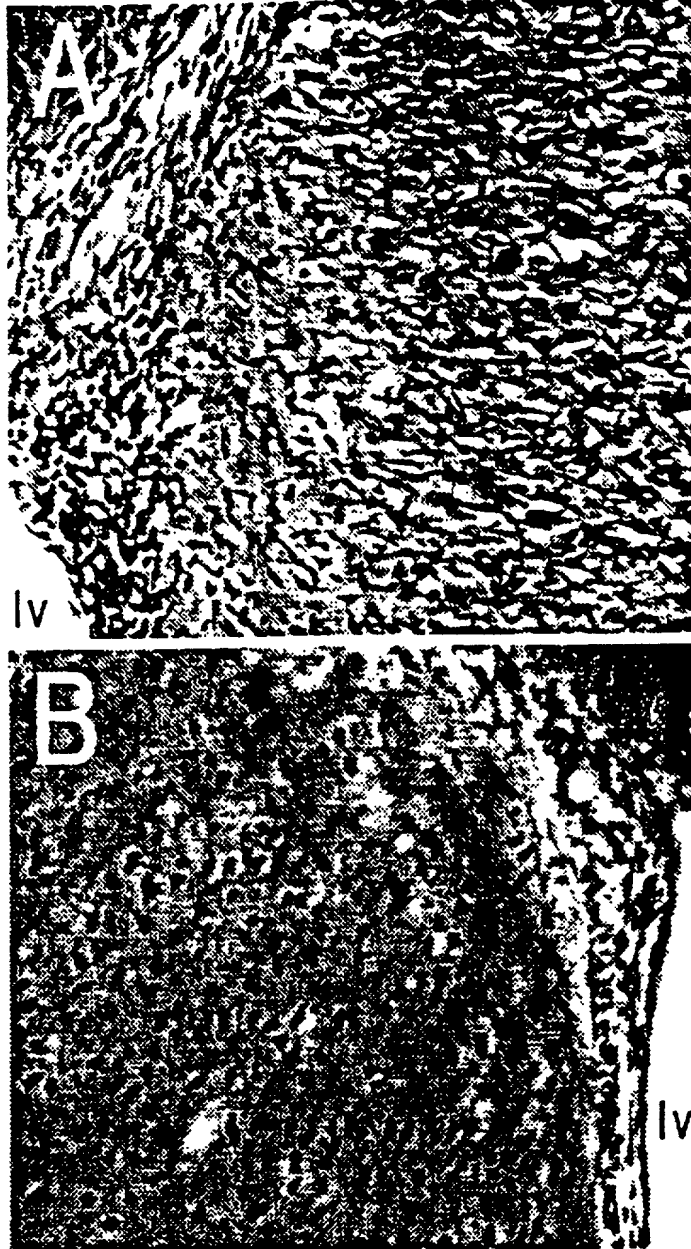


Fig. 6

Bar graph showing the number of dendritic profiles per mm² (d p.m./mm²) in the ridge, striatum, and SEZ for ipsilateral (white bars) and contralateral (hatched bars) sides. The y-axis ranges from 0 to 900. The x-axis categories are ridge, striatum, and SEZ. Error bars are shown for each bar. Significance markers (* and **) are present above the bars.

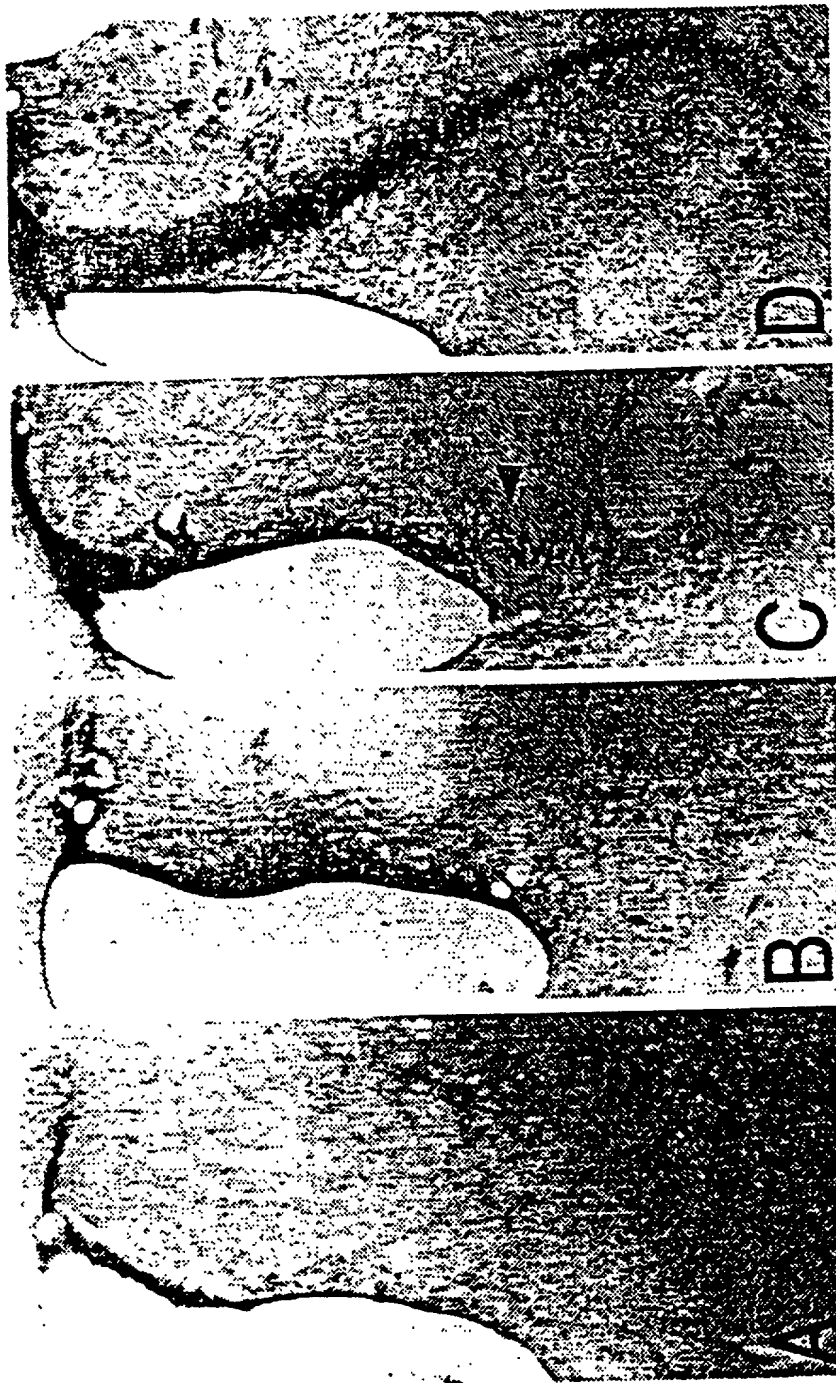
Region	Ipsi (d p.m./mm²)	Contra (d p.m./mm²)	Significance
ridge	~770	~770	
striatum	~160	~170	*
SEZ	~690	~440	**

Fig. 7



Figs. 8A-8B

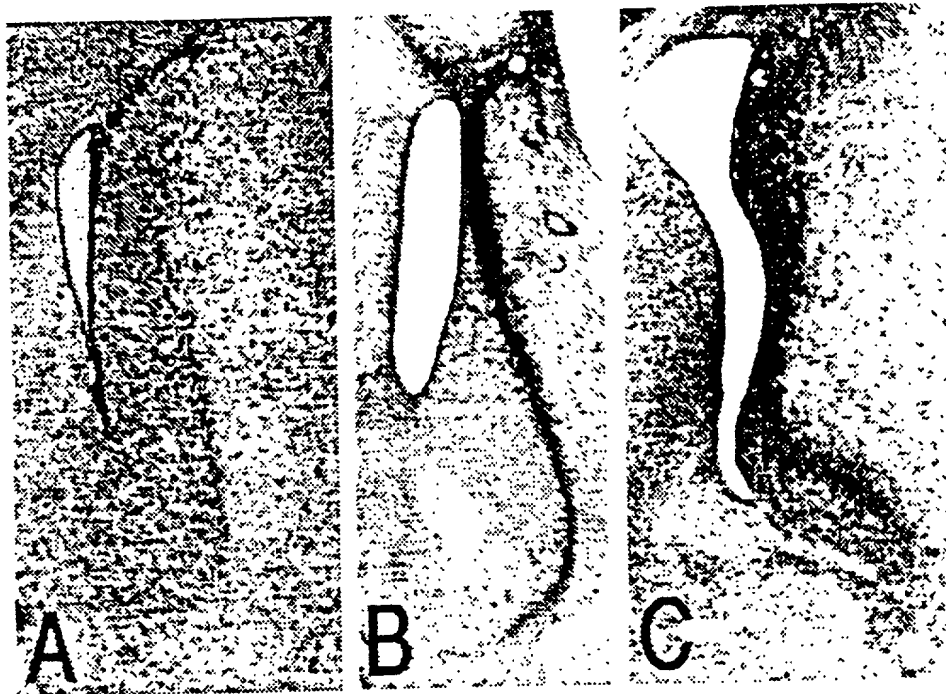
008727" E66E260



Figs. 9A-9D

The figure consists of two electron micrographs, labeled A and B, showing the ultrastructure of a cell. Both images show a cell with a prominent nucleus and a large, electron-lucent area. Image A shows a cell with a more electron-dense nucleus and a large, electron-dense area. Image B shows a cell with a more electron-dense nucleus and a large, electron-dense area.

Figs. 10A-10B



Figs. 11A-11C

003727 4666260

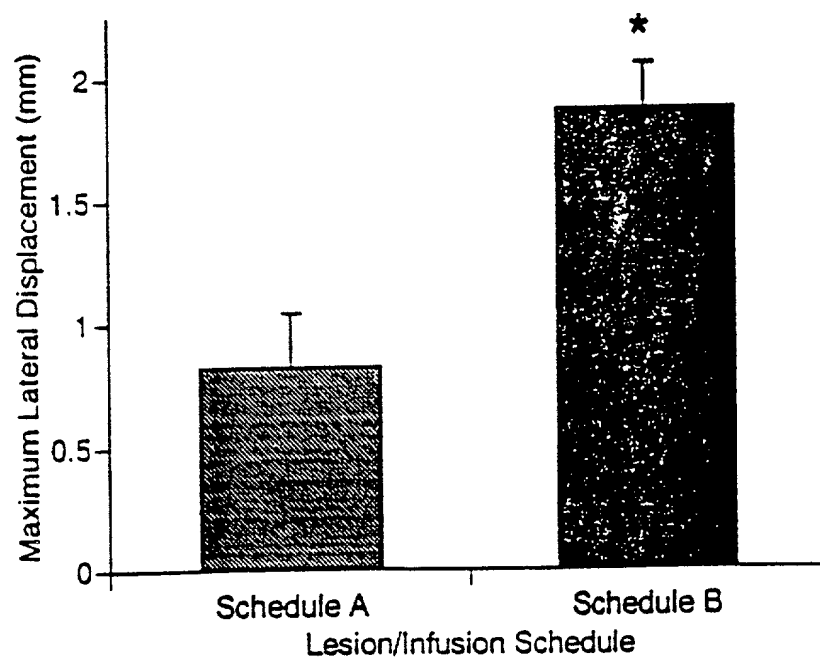


Fig. 12